

XX Claim 3; Page 640; 837pp; English.
PS
XX
CC 242265 to 243075 represent novel 5' expressed sequence tag (EST)
CC sequences, corresponding to human secreted proteins. Y64651 to Y65438
CC represent the EST-related proteins corresponding to 242265 to 243052.
CC The 5' ESTs can be used for producing secreted human gene products.
CC They can be used to identify and isolate 5' untranslated regions (UTRs)
CC and upstream regulatory regions which control the location, development
CC stage, rate, and quantity of protein synthesis, as well as stability of
CC mRNA. The ESTs are also useful as probes for chromosome mapping, and to
CC obtain full length cDNA clones. The ESTs can also be used in forensic
CC procedures to identify individuals, or in diagnostic procedures to
CC identify individuals having genetic diseases resulting from abnormal
CC gene expression. The products may also be used in gene therapy protocols.
CC The nucleic acids encoding signal peptides can be used for directing
CC extracellular secretion of a polypeptide or the insertion of a
CC polypeptide into a membrane, or importing a polypeptide into a cell.
CC The proteins encoded by the EST sequences may be useful in treating a
CC variety of human conditions. Secreted proteins have therapeutic value,
CC and the identification of new secreted proteins is valuable. 242249 to
CC 242264 and Y64644 to Y64650 represent sequences used in the
CC exemplification of the present invention.
SQ Sequence 66 AA;

Query Match 100.0%; Score 77; DB 21; Length 66;
Best Local Similarity 20.7%; Pred. No. 20;
Matches 6; Conservative 23; Mismatches 0; Indels 0; Gaps 0;

QY 1 CXXCXXCXXCXXCXXCXXCXXCXXCXXC 29
|::|::|::|::|::|::|::|::|::|::|
Db 27 cpcvcmccvcmccvcmccvcmccvcmccv 55

RESULT 2

W62830 ID W62830 standard; Protein: 625 AA.

XX AC W62830;

XX DF 27-OCT-1998 (first entry)

XX DE Macadamia integrifolia antimicrobial protein.

XX KM antimicrobial protein; infestation; control.

XX OS Macadamia integrifolia.

XX FH Key Location/Qualifiers

FT Peptide 1..28

FT Protein /note="signal peptide"

FT Protein /note="mature protein"

XX WO9827805-A1.

XX PN 02-JUL-1998.

XX PD 22-DEC-1997; 97WO-AU00874.

XX PR 20-DEC-1996; 96AU-0004275.

XX PA (RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY.

XX PI Bower NT, Goulter KC, Green JL, Manners JM, Marcus JP;

XX DR WPI: 1998-377279/32.

XX DR N-PSDB; V42316.
XX PT Novel anti-microbial protein from e.g. Macadamia integrifolia -
XX useful for controlling microbial infestations of plants or mammals

XX Claim 1; Page 43-45; 96pp; English.
PS
XX
CC The sequence is that of an antimicrobial protein which can
CC be used to control microbial infestations in plants and mammalian
CC animals.
XX
SQ Sequence 625 AA;

Query Match 100.0%; Score 77; DB 19; Length 625;
Best Local Similarity 20.7%; Pred. No. 1.8e+02;
Matches 6; Conservative 23; Mismatches 0; Indels 0; Gaps 0;

QY 1 CXXCXXCXXCXXCXXCXXCXXCXXCXXC 29
|::|::|::|::|::|::|::|::|::|::|
Db 41 cqqcqrccrccqesprqgycqrcckeic 69

RESULT 3

W62828 ID W62828 standard; Protein: 666 AA.

XX AC W62828;

XX DT 27-OCT-1998 (first entry)

XX DE Macadamia integrifolia antimicrobial protein.

XX KM antimicrobial protein; infestation; control.

XX OS Macadamia integrifolia.

XX FH Key Location/Qualifiers

FT Peptide 1..28

FT Protein /note="signal peptide"

FT Protein /note="mature protein"

XX WO9827805-A1.

XX PN 02-JUL-1998.

XX PD 22-DEC-1997; 97WO-AU00874.

XX PR 20-DEC-1996; 96AU-0004275.

XX PA (RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY.

XX PI Bower NT, Goulter KC, Green JL, Manners JM, Marcus JP;

XX DR WPI: 1998-377279/32.

XX DR N-PSDB; V42310.

XX PT Novel anti-microbial protein from e.g. Macadamia integrifolia -
XX useful for controlling microbial infestations of plants or mammals

XX PS Claim 1; Page 34-36; 96pp; English.

XX CC The sequence is that of an antimicrobial protein which can
XX be used to control microbial infestations in plants and mammalian
XX animals.

XX SQ Sequence 666 AA;

Query Match 100.0%; Score 77; DB 19; Length 666;
Best Local Similarity 20.7%; Pred. No. 2e+02;
Matches 6; Conservative 23; Mismatches 0; Indels 0; Gaps 0;

QY 1 CXXCXXCXXCXXCXXCXXCXXCXXCXXC 29
|::|::|::|::|::|::|::|::|::|::|
Db 82 cqqcqrccrccqesprqgycqrcckeic 110

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RESULT 4
W62829 ID W62829 standard; Protein; 666 AA.
XX
AC W62829;
XX
XX 27-OCT-1998 (first entry)
XX
DE Macadamia integrifolia antimicrobial protein.
XX
XX antimicrobial protein; infestation; control.
XX
KM Macadamia integrifolia.
XX
OS
XX
XX Key Location/Qualifiers
FH Peptide 1..28
FT /note="signal peptide"
FT Protein 29..666
FT /note="mature protein"
XX
XX W09827805-A1.
XX
XX 02-JUL-1998.
XX
XX 22-DEC-1997; 97MO-AU00874.
XX
XX 20-DEC-1996; 96AU-0004275.
XX
XX (RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY.
XX
XX Bower NI, Goulter KC, Green JL, Manners JM, Marcus JP;
XX
XX WPI: 1998-377279/32.
XX
XX N-PSDB; V42311.
XX
XX Novel anti-microbial protein from e.g. Macadamia integrifolia -
XX
XX useful for controlling microbial infestations of plants or mammals
XX
XX Claim 1; Page 39-41; 96pp; English.
XX
XX The sequence is that of an antimicrobial protein which can
XX
XX be used to control microbial infestations in plants and mammalian
XX
XX animals.
XX
XX Sequence 666 AA:
SQ
Query Match 100.0%; Score 77; DB 19; Length 666;
Best Local Similarity 20.7%; Pred. No. 2e+02;
Matches 6; Conservative 23; Mismatches 0; Indels 0; Gaps 0;
OY 1 CXXCXXCXXCXXCXXCXXCXXCXXCXXC 29
DB 82 cqgcqrrcrqgsqpsrqsycqrrckeic 110
RESULT 5
Y70731 ID Y70731 standard; protein; 31 AA.
XX
XX Y70731;
XX
XX 24-JUL-2000 (first entry)
XX
XX Wnt antagonist protein consensus sequence-1.
XX
XX Wnt antagonist; contraceptive; contraceptive vaccine; oocyte development;
XX
XX female primate contraception; oocyte viability.
XX
XX Synthetic.
XX
XX

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FH Key Location/Qualifiers
FT Misc-difference 2
FT /label= Unknown
FT /note="Xaa may be 9 amino acids in length; some
FT amino acids may be absent"
FT Misc-difference 4
FT /label= Unknown
FT /note="Xaa may be 42 amino acids in length; some
FT amino acids may be absent"
FT Misc-difference 14
FT /label= Unknown
FT Misc-difference 15
FT /label= Unknown
FT Misc-difference 16
FT /label= Unknown
FT Misc-difference 17
FT /label= Unknown
FT Misc-difference 18
FT /label= Unknown
FT Misc-difference 19
FT /label= Unknown
FT Misc-difference 21
FT /label= Unknown
FT /note="Xaa may be 10 amino acids in length; some
FT amino acids may be absent"
FT Misc-difference 23
FT /label= Unknown
FT Misc-difference 24
FT /label= Unknown
FT Misc-difference 25
FT /label= Unknown
FT Misc-difference 27
FT /label= Unknown
FT /note="Xaa may be 7 amino acids in length; some
FT amino acids may be absent"
FT Misc-difference 29
FT /label= Unknown
FT /note="Xaa may be 27 amino acids in length; some
FT amino acids may be absent"
FT Misc-difference 31
FT /label= Unknown
FT /note="Xaa may be 13 amino acids in length; some
FT amino acids may be absent"
FH W0200021555-A1.
PD 20-APR-2000.
XX
XX 13-OCT-1999; 99MO-US23640.
XX
XX 15-OCT-1998; 98US-0104355.
XX
XX (HARD ) HARVARD COLLEGE.
XX
XX McMahon AP, Parr BA, Vaino S;
XX
XX WPI: 2000-317845/27.
XX
XX Contraceptive composition for inhibiting oocyte development in a female
XX primate comprises a Wnt polypeptide antagonist -
XX
XX Claim 12; Page 44; 57pp; English.
XX
XX The patent discloses a method of female primate contraception comprising
XX administering an antagonist of a Wnt polypeptide, inhibiting oocyte
XX development. Wnt polypeptides are useful for promotive maturation of an
XX immature oocyte. Wnt polypeptides are also useful for increasing the
XX number of mature oocytes and to enhance oocyte viability. The present
XX peptide is a consensus sequence of Wnt antagonist which inhibits the
XX physiological activity of a Wnt polypeptide. Antagonistic polypeptides
XX may contain a cysteine-rich domain.
XX
XX Sequence 31 AA:
SQ

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XX AC Y35935;
XX DT 13-SEP-1999 (first entry)
XX DE Extended human secreted protein sequence, SEQ ID NO. 184.
XX KM Secreted protein; human; cytokine; cellular proliferation; cell movement;
XX KM cellular differentiation; immune system regulator; anti-inflammatory;
XX KM haematopoiesis regulator; tissue growth regulator; tumour inhibitor;
XX KM reproductive hormone regulator; chemotaxis; chemokinesis; gene therapy;
XX OS genetic disease.
XX OS Homo sapiens.
XX PN MO9931236-A2.
XX PD 24-JUN-1999.
XX PF 17-DEC-1998; 98WO-1B02122.
XX PR 10-AUG-1998; 98US-0096116.
XX PR 17-DEC-1997; 97US-0069957.
XX PR 09-FEB-1998; 98US-0074121.
XX PR 13-APR-1998; 98US-0081563.
XX PA (GEST ) GENSET.
XX PI Bougueleret L, Duclert A, Dumas Milne Edwards J;
XX DR WPI; 1999-385906/32.
XX DR N-PSDB; X97619.
XX PT New isolated human secreted proteins
XX PS Claim 9; Page 215-216; 516pp; English.
XX XX This sequence is encoded by an extended human secreted protein coding
XX CC sequence of the invention. The secreted proteins can be used in treating
XX CC or controlling a variety of human conditions. The secreted proteins may
XX CC act as cytokines or may affect cellular proliferation or differentiation
XX CC or may act as immune system regulators, haematopoiesis regulators, tissue
XX CC growth regulators, regulators of reproductive hormones or cell movement
XX CC or have chemotactic/chemokinetic, receptor/ligand, anti-inflammatory or
XX CC tumour inhibition activity. The DNAs can be used in forensic procedures
XX CC to identify individuals or in diagnostic procedures to identify
XX CC individuals having genetic diseases resulting from abnormal expression of
XX CC the genes corresponding to the extended cDNAs. They are also useful for
XX CC constructing a high resolution map of the human chromosomes. They can
XX CC also be used for gene therapy to control or treat genetic diseases.
XX SQ Sequence 73 AA:
XX
Query Match 88.3%; Score 68; DB 20; Length 73;
Best Local Similarity 17.9%; Pred. No. 1e+02;
Matches 5; Conservative 23; Mismatches 0; Indels 0; Gaps 0;
QY 2 XXCXXXCXXXXXXXXXXXXCXXCXXC 29
DB 42 dlcjhscsfqkccetnkiccatcgnic 69
XX
RESULT 9
Y68907
XX ID Y68907 standard; Protein; 76 AA.
XX AC Y68907;
XX DT 16-MAY-2000 (first entry)
XX DE A mouse WDMN-2 protein (homologous to TANGO-175).
XX

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XX KM TANGO-175; proteinase; metastasis; inflammation; cancer; inflammation;
XX KM bone marrow accessory cell; hematopoietic stem cell growth;
XX KM hematopoietic disorder; genetic lesion; WDMN-2.
XX OS Mus sp.
XX PN MO200006699-A1.
XX PD 10-FEB-2000.
XX PF 29-JUL-1999; 99WO-US17289.
XX PR 29-JUL-1998; 98US-0124538.
XX PA (MILL-) MILLENIUM BIOTHERAPEUTICS INC.
XX PI McCarthy SA;
XX DR WPI; 2000-183117/16.
XX DR N-PSDB; Z60804.
XX PT Novel human and murine TANGO-175 and murine WDMN-2 nucleic acids and
XX PT proteins useful for treatment and diagnosis of cancer, inflammation and
XX PT hematopoietic disorders
XX PS Example 3; Fig 3; 134pp; English.
XX CC The present sequence represents mouse WDMN-2, which has homology to
XX CC TANGO-175. TANGO-175 is related to several proteins in the four
XX CC disulphide core family. Human TANGO-175 has activities similar to that
XX CC of anti-leukoproteinase and WDMN-1 and may therefore have a role similar
XX CC to these proteins by inhibiting proteinases associated with metastasis.
XX CC The protein may play a role in regulating inflammation and also in the
XX CC growth of hematopoietic stem cells by neutralizing proteinases produced
XX CC by bone marrow accessory cells. TANGO-175 is therefore useful in
XX CC treatment and diagnosis of cancer, inflammation and hematopoietic
XX CC disorders. Primers and probes, which hybridize to human TANGO-175
XX CC nucleic acid molecules and antibodies against human TANGO-175 protein,
XX CC are useful for detecting the presence of the nucleic acid molecule or
XX CC protein in a sample. The proteins and nucleic acids can be used to screen
XX CC drugs or compounds, which modulate TANGO-175 activity or expression, to
XX CC detect genetic lesions and to modulate TANGO-175 activity.
XX SQ Sequence 76 AA:
XX
Query Match 88.3%; Score 68; DB 21; Length 76;
Best Local Similarity 17.9%; Pred. No. 1e+02;
Matches 5; Conservative 23; Mismatches 0; Indels 0; Gaps 0;
QY 2 XXCXXXCXXXXXXXXXXXXCXXCXXC 29
DB 44 gicvdqsgdsgcpymkcsnscghvc 71
XX
RESULT 10
Y69209
XX ID Y69209 standard; Protein; 92 AA.
XX AC Y69209;
XX DT 30-MAY-2000 (first entry)
XX DE Amino acid sequence of honey bee venom PK3.101 protein.
XX XX Protein PK3.101; honey bee; venom; interleukin-8; IL-8; receptor;
XX KM CXCR1; CXCR2; cyclooxygenase; lipoxigenase; phospholipase; protease;
XX KM inflammatory disease; gene therapy; cancer; autoimmune disease; pain;
XX KM chemokine imbalance; rheumatoid arthritis; multiple sclerosis;
XX KM psoriasis; systemic lupus erythematosus; Crohn's disease; vasculitis;
XX KM scleroderma; metastatic cancer; Alzheimer's disease; wound healing;
XX KM aging process; antigen.
XX

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OS  Apis mellifera.
XX
FH  Key      Location/Qualifiers
FT  Peptide  1..19
FT          /note= "Signal peptide"
FT  Region   20..34
FT          /note= "this region contains 5 GGX repeats"
XX
XX  GB2341389-A.
XX
XX  15-MAR-2000.
XX
XX  13-SEP-1999; 99GB-0021605.
XX
XX  14-SEP-1998; 98US-0100172.
XX
XX  (PANP-) PAN PACIFIC PHARM INC.
XX
XX  Chi X, Lu Y;
XX
XX  WPI: 2000-185368/17.
XX  N-PSDB; 261247.
XX
XX  Isolated nucleic acids encoding the bee venom protein PX3.101 useful
PT  for treating autoimmune and inflammatory disorders such as rheumatoid
PT  arthritis.
XX
XX  Claim 2: Flg 3A-B: 83pp; English.
XX
XX  The present sequence represents the protein PX3.101, which is a honey
CC  bee venom isolated Apis mellifera. PX3.101 inhibits the binding of
CC  Interleukin-8 (IL-8) to its receptor (e.g. CXCR1 and CXCR2) and
CC  inhibits a variety of enzymes (e.g. cyclooxygenases, lipoxigenases,
CC  phospholipases and proteases) associated with inflammatory diseases.
CC  The nucleic acids may be used for the recombinant production of
CC  PX3.101 proteins either in vivo (as part of a gene therapy protocol)
CC  or in vitro (as a fermentation culture). The nucleic acids may also
CC  be used as probes to identify similar sequences in samples. The PX3.101
CC  protein may be used for the treatment of inflammatory diseases, cancers,
CC  autoimmune diseases, pain and/or diseases associated with chemokine
CC  (especially IL-8) imbalances such as rheumatoid arthritis, multiple
CC  sclerosis, psoriasis, systemic lupus erythematosus (SLE), Crohn's
CC  disease, vasculitis, scleroderma, metastatic cancer and Alzheimer's
CC  disease in humans. It is also disclosed that the proteins may be used
CC  to accelerate wound healing, reduce several aging processes and protect
CC  against ultraviolet light. The proteins may also be used as antigens in
CC  the production of antibodies specific for PX3.101. The antibodies may
CC  be used as diagnostic agents to detect PX3.101 protein in samples and
CC  to down regulate PX3.101 activity.
XX
XX  Sequence 92 AA:
SQ

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Query Match 88.3%; Score 68; DB 21; Length 92;
 Best Local Similarity 17.9%; Pred. No. 1.3e+02;
 Matches 5; Conservative 23; Mismatches 0; Indels 0; Gaps 0;

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OY  2 XXXXXXXXXXXXXXXXXXXXCCXXC 29
    ::::::::::::::::::::::|
DB  44 srodcgrcfcpnvvpkplckitcapgc 71

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RESULT 11
 Y36164
 ID Y36164 standard; protein: 93 AA.
 AC Y36164;
 XX
 DT 23-SEP-1999 (first entry)
 XX
 DE Human secreted protein #36.
 XX
 KW Secreted protein; human; cytosstatic; thrombotic; osteopathic; forensic;

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KW  diagnostic; gene therapy; chromosome mapping; secretion vector.
XX
XX  Homo sapiens.
XX
XX  MO9925825-A2.
XX
XX  27-MAY-1999.
XX
XX  13-NOV-1998; 98WO-IB01862.
XX
XX  04-SEP-1998; 98US-0099273.
XX  13-NOV-1997; 97US-0066677.
XX  17-DEC-1997; 97US-0069957.
XX  09-FEB-1998; 98US-0074121.
XX  13-APR-1998; 98US-0081563.
XX  10-AUG-1998; 98US-0096116.
XX
XX  (GEST ) GENSET.
XX
XX  Bougueleret L, Duclet A, Dumas Milne Edwards J;
XX
XX  WPI: 1999-347472/29.
XX  N-PSDB; X97848.
XX
XX  Extended cDNAs encoding secreted proteins
XX
XX  Example 28: Page 246; 307pp; English.
XX
XX  Y36129-Y36222 represent novel human secreted proteins encoded by the
CC  extended cDNA sequences represented in X97813-X97906. The proteins
CC  of the invention have cytosstatic, thrombotic and osteopathic activity.
CC  The extended cDNAs can be used to express secreted proteins or parts of
CC  them or to obtain antibodies capable of binding to the secreted proteins.
CC  They may also be used in diagnostic, forensic, gene therapy and
CC  chromosome mapping procedures. Uses also include design of expression
CC  vectors and secretion vectors.
XX
XX  Sequence 93 AA:
SQ

```

Query Match 88.3%; Score 68; DB 20; Length 93;
 Best Local Similarity 17.9%; Pred. No. 1.3e+02;
 Matches 5; Conservative 23; Mismatches 0; Indels 0; Gaps 0;

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OY  2 XXXXXXXXXXXXXXXXXXXXCCXXC 29
    ::::::::::::::::::::::|
DB  43 encthtctmgedckgfgccstfcjvc 70

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RESULT 12
 Y36211
 ID Y36211 standard; protein: 93 AA.
 AC Y36211;
 XX
 DT 23-SEP-1999 (first entry)
 XX
 DE Human secreted protein #83.
 XX
 KW Secreted protein; human; cytosstatic; thrombotic; osteopathic; forensic;
 KW diagnostic; gene therapy; chromosome mapping; secretion vector.
 XX
 OS Homo sapiens.
 XX
 PN MO9925825-A2.
 XX
 PD 27-MAY-1999.
 XX
 PF 13-NOV-1998; 98WO-IB01862.
 XX
 PR 04-SEP-1998; 98US-0099273.
 PR 13-NOV-1997; 97US-0066677.
 PR 17-DEC-1997; 97US-0069957.
 PR 09-FEB-1998; 98US-0074121.
 PR 13-APR-1998; 98US-0081563.
 PR 10-AUG-1998; 98US-0096116.
 PR
 PR 17-DEC-1997; 97US-0069957.

PS Disclosure: Page 16-17; 28pp; English.
 XX
 CC DNA sequences have been identified by constructing a cDNA library
 CC from human epididymis RNA, screening with epididymis and testis
 CC probes, and further screening with brain and liver probes.
 CC The DNA mol. is useful in the cloning and expression of human
 CC epididymis-specific polypeptides in pro- or eukaryotic host cells.
 CC The polypeptides and antibodies to the peptides are useful for
 CC diagnosis or therapy of male infertility and for
 CC immunosterilisation of mammals.
 XX
 SQ Sequence 125 AA:

Query Match 88.3%; Score 68; DB 12; Length 125;
 Best local Similarity 17.9%; Pred. No. 1.7e+02;
 Matches 5; Conservative 23; Mismatches 0; Indels 0; Gaps 0;
 QY 2 XXCXCCXXXXXXXXXXCCXCCXCC 29
 :|::|::|::|::|::|::|::|::|::|
 Db 43 qnctgecvdsccadnlkccsagcatfc 70

RESULT 15

W81779
 ID W81779 standard; Protein; 125 AA.

AC W81779;

DT 23-FEB-1999 (first entry)

DE Human HE4 protein.

KW HE4; epididymis-specific; diagnosis; male infertility; treatment;
 KW sterility; immunosterilisation.

OS Homo sapiens.

Key Location/Qualifiers
 FT Peptide 1..30
 FT /label= signal
 FT Protein 31..125
 FT /label= HE4

EP878544-A1.

18-NOV-1998.

29-JAN-1991; 91EP-0250021.

30-NOV-1990; 90DE-4038189.

01-FEB-1990; 90DE-4002981.

(IHFF-) IHF INST HORMON & FORTPFLANZUNGS.

Ivell R, Kirchhoff C;

WPI: 1998-585748/50.

N-PSDB; V64622.

DNA encoding human epididymis polypeptides - useful for, e.g.
 diagnosis of male infertility

Disclosure: Page 16-17; 29pp; German.

This sequence represents a novel human epididymis-specific protein, HE4.
 This sequence may be used for cloning and for expression of human
 epididymis-specific polypeptides in prokaryotic or eukaryotic host
 cells. Such proteins and antibodies generated from them may be used for
 diagnosis of e.g. male infertility. The polypeptides and antibodies
 may also be used for treatment of male infertility and for
 immunosterilisation of mammals.

SQ Sequence 125 AA:

Query Match 88.3%; Score 68; DB 19; Length 125;
 Best local Similarity 17.9%; Pred. No. 1.7e+02;
 Matches 5; Conservative 23; Mismatches 0; Indels 0; Gaps 0;
 QY 2 XXCXCCXXXXXXXXXXCCXCCXCC 29
 :|::|::|::|::|::|::|::|::|::|
 Db 43 qnctgecvdsccadnlkccsagcatfc 70

Search completed: March 1, 2001, 16:18:27
 Job time: 496 sec